Revised 09/09/2003

CRF Errors Edited by the STIC Systems Branch

Seria	Realigned nucleic acid/amino acid numbers/text text "wrapped" to the next line	CRF Edit Date: 3/8/04 Edited by:
	Corrected the SEQ ID NO. Sequence numbers of	edited were:
ran ra ur	Inserted or corrected a nucleic number at the end NO's edited:	d of a nucleic line. SEQ ID
	Deleted: invalid beginning/end-of-file text;_	page numbers
	Inserted mandatory headings/numeric identifiers	, specifically:
	Moved responses to same line as heading/numeric	c identifier, specifically:
<u>J</u>	Other: Sequerce 4-concerted asserts as	id rumbering



IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/785,673

DATE: 03/08/2004 TIME: 15:22:42

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\03082004\J785673.raw

SEQUENCE LISTING

```
4 (1) GENERAL INFORMATION:
              (i) APPLICANT: MINETTI, CONCEICAO;
       6
                             MICHON, FRANCIS;
      8
                             PULLEN, JEFFREY K.;
      9
                             POLDVINO-BODNAR, MARYELLEN;
      10
                             LIANG, SHU-MEI;
     11
                             TAI, JOSEPH Y.
     13
             (ii) TITLE OF INVENTION: MODIFIED IMMUNOGENIC
     14
                                       PNEUMOLYSIN COMPOSITIONS AS VACCINES
     16
            (iii) NUMBER OF SEQUENCES: 18
     18
             (iv) CORRESPONDENCE ADDRESS:
     19
                   (A) ADDRESSEE: MORGAN & FINNEGAN, L.L.P.
     20
                   (B) STREET: 345 PARK AVENUE
     21
                   (C) CITY: NEW YORK
     22
                   (D) STATE: NEW YORK
     23
                   (E) COUNTRY: USA
     24
                   (F) ZIP: 10154
     26
             (v) COMPUTER READABLE FORM:
     27
                   (A) MEDIUM TYPE: FLOPPY DISK
     28
                   (B) COMPUTER: IBM PC COMPATIBLE
     29
                   (C) OPERATING SYSTEM: PC-DOS/MS-DOS
     30
                   (D) SOFTWARE: MICROSOFT WORD 97
            (vi) CURRENT APPLICATION DATA:
     32
C--> 33
                   (A) APPLICATION NUMBER: US/10/785,673
C--> 34
                   (B) FILING DATE: 23-Feb-2004
     35
                  (C) CLASSIFICATION:
     41
           (vii) PRIOR APPLICATION DATA:
     38
                  (A) APPLICATION NUMBER: 60/053,306
    39
                  (B) FILING DATE: 1997-07-21
    42
                  (A) APPLICATION NUMBER: 60/073,456
    43
                  (B) FILING DATE: 1998-02-02
    45
          (viii) ATTORNEY/AGENT INFORMATION:
    46
                  (A) NAME: DARRYL H. STEENSMA
    47
                  (B) REGISTRATION NUMBER: 43,155
    49
                  (C) REFERENCE/DOCKET NUMBER: 1758-4036US2
    51
            (ix) TELECOMMUNICATION INFORMATION:
    52
                  (A) TELEPHONE: (212) 758-4800
    53
                  (B) TELEFAX: (212) 751-6849
                  (C) TELEX: 421792
    57 (2) INFORMATION FOR SEQ ID NO: 1:
    59
            (i) SEQUENCE CHARACTERISTICS:
```

(A) LENGTH: 1413

60

DATE: 03/08/2004

TIME: 15:22:42

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/785,673

Input Set : A:\PTO.AMC.txt

```
61
                  (B) TYPE: nucleic acid
                  (C) STRANDEDNESS: double
                  (D) TOPOLOGY: linear
            (ii) MOLECULE TYPE: cDNA
            (vi) ORIGINAL SOURCE:
     68
                  (A) ORGANISM: S. pneumoniae
     70
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
C--> 72 ATGGCAAATA AAGCAGTAAA TGACTTTATA CTAGCTATGA 40
     74 ATTACGATAA AAAGAAACTC TTGACCCATC AGGGAGAAAG 80
    76 TATTGAAAAT CGTTTCATCA AAGAGGGTAA TCAGCTACCC 120
    78 GATGAGTTTG TTGTTATCGA AAGAAAGAAG CGGAGCTTGT 160
    80 CGACAAATAC AAGTGATATT TCTGTAACAG CTACCAACGA 200
    82 CAGTCGCCTC TATCCTGGAG CACTTCTCGT AGTGGATGAG 240
    84 ACCTTGTTAG AGAATAATCC CACTCTTCTT GCGGTCGATC 280
    86 GTGCTCCGAT GACTTATAGT ATTGATTTGC CTGGTTTGGC 320
    88 AAGTAGCGAT AGCTTTCTCC AAGTGGAAGA TCCCAGCAAT 360
    90 TCAAGTGTTC GCGGAGCGGT AAACGATTTG TTGGCTAAGT 400
    92 GGCATCAAGA TTATGGTCAG GTCAATAATG TCCCAGCTAG 440
    94 AATGCAGTAT GAAAAAATCA CGGCTCACAG CATGGAACAA 480
    97 CTCAAGGTCA AGTTTGGTTC TGACTTTGAA AAGACAGGGA 520
    99 ATTCTCTTGA TATTGATTTT AACTCTGTCC ATTCAGGCGA 560
    101 AAAGCAGATT CAGATTGTTA ATTTTAAGCA GATTTATTAT 600
    103 ACAGTCAGCG TAGACGCTGT TAAAAATCCA GGAGATGTGT 640
    105 TTCAAGATAC TGTAACGGTA GAGGATTTAA AACAGAGAGG
    107 AATTTCTGCA GAGCGTCCTT TGGTCTATAT TTCGAGTGTT
    109 GCTTATGGGC GCCAAGTCTA TCTCAAGTTG GAAACCACGA
    111 GTAAGAGTGA TGAAGTAGAG GCTGCTTTTG AAGCTTTGAT 800
    113 AAAAGGAGTC AAGGTAGCTC CTCAGACAGA GTGGAAGCAG 840
    115 ATTTTGGACA ATACAGAAGT GAAGGCGGTT ATTTTAGGGG 880
    117 GCGACCCAAG TTCGGGTGCC CGAGTTGTAA CAGGCAAGGT 920
    119 GGATATGGTA GAGGACTTGA TTCAAGAAGG CAGTCGCTTT 960
    121 ACAGCAGATC ATCCAGGCTT GCCGATTTCC TATACAACTT 1000
    123 CTTTTTTACG TGACAATGTA GTTGCGACCT TTCAAAATAG 1040
    125 TACAGACTAT GTTGAGACTA AGGTTACAGC TTACAGAAAC 1080
   127 GGAGATTTAC TGCTGGATCA TAGTGGTGCC TATGTTGCCC 1120
   129 AATATTATAT TACTTGGAAT GAATTATCCT ATGATCATCA 1160
   131 AGGTAAGGAA GTCTTGACTC CTAAGGCTTG GGACAGAAAT 1200
   133 GGGCAGGATT TAACGGCTCA CTTTACCACT AGTATTCCTT 1240
   135 TAAAAGGGAA TGTTCGTAAT CTCTCTGTCA AAATTAGAGA 1280
   137 GTGTACCGGG CTTGCTTGGG AATGGTGGCG TACGGTTTAT 1320
   139 GAAAAAACCG ATTTGCCACT AGTGCGTAAG CGGACGATTT 1360
   141 CTATTTGGGG AACAACTCTC TATCCGCAGG TAGAAGATAA 1400
   144 GGTAGAAAAT GAC 1413
   147 (2) INFORMATION FOR SEQ ID NO: 2:
             (i) SEQUENCE CHARACTERISTICS:
   149
   150
                  (A) LENGTH: 1413
   151
                  (B) TYPE: nucleic acid
   152
                  (C) STRANDEDNESS: double
   153
                 (D) TOPOLOGY: linear
```

PATENT APPLICATION: US/10/785,673

DATE: 03/08/2004 TIME: 15:22:42

Input Set : A:\PTO.AMC.txt

```
155
               (ii) MOLECULE TYPE: cDNA
       157
               (vi) ORIGINAL SOURCE:
       158
                     (A) ORGANISM: S. pneumoniae
      160
               (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
 C--> 162 ATGGCAAATA AAGCAGTAAA TGACTTTATA CTAGCTATGA 40
      164 ATTACGATAN AAANAAACTC TTGACCCATC AGGGAGAAAG 80
      166 TATTGAAAAT CGTTTCANCA AAGAGGGTAA TCAGCTACCC 120
      168 GNTGAGTTTG TTGNTANCGA AAGAAAGAAG CGGAGCTTGT 160
      170 CGACAAATAC AAGTGATATT NCTGTANCAG CTACCNACGA 200
      172 CAGTCGCCTC TATCCTGGAG CACTTCTCGT AGTGGATGAG 240
      174 ACCTTGTNAG AGAATAATCC CACTCTTCTT GCGGTNGATC 280
      176 GTGCTCCGAT GACTTATAGT ANTGNTTTGC CTGGTTTGGC 320
      178 AAGTAGCGAT AGCTTTCTCC AAGTGGAAGA NCCCAGCAAT 360
      180 TCAAGTGTTC GCGGAGCGGN ANACGATTTG TTGGCTAAGT 400
      182 GGCATCAAGA TTATGGTCAG GTCAATAATG TCCCAGCTAG 440
      184 AANGCAGTAT GAAAAAATNA CGGCTCACAG CATGGAACAA 480
      186 CTCAAGGTCA AGTTTGGTTC TGACTTTGAA AAGNCAGGGA 520
      188 ATTCTCTTGA TATTGATTTT AACTCTGTCC ATTCAGGNGA 560
      191 AAAGCNGATT CAGATTGTTA ATNTTAAGCA GATTTATTAT 600
      193 ACAGTCAGCG TAGACGCTGT TAAAAATCCA GGAGATGTGT 640
      195 TTCAAGATAC TGTAACGGTA GAGGATTTAA AACAGAGAG 680
      197 AATTTCTGCA GAGCGTCCTT TGGTCTATAT TTCGAGNGTT
      199 GCTTATGGGC GCCAAGTCTA TCTCAAGTTG GAAACCACGA 760
      201 GTANGAGTGN TGAAGTAGAG GCTGCTTTTG AAGCTTTGAT 800
      203 AAAAGGAGTC AAGGTAGCTC CTCAGACAGA GTGGAAGCAG
      205 ATTTTGGACA ATACAGAAGT GAAGGCGGTT ATTTTAGGGG 880
      207 GCGACCCAAG TTCGGGTGCC CGAGTTGTAA CAGGCAAGGT 920
     209 GGATATGGTA GAGGACTTGA TTCAAGAAGG CAGTCGCTTT 960
     211 ACAGCAGATC ATCCAGGCTT GCCGATTTCC TATACAACTT 1000
     213 CTTTTTTACG TGACAATGTA GTTGCGACCT TTCAAAANAG 1040
     215 TACAGACTAT GTTGAGACTA AGGTTACAGC TTACAGAAAC 1080
     217 GGAGATTTAC TGCTGGATCA TAGTGGTGCC TATGTTGCCC 1120
     219 AATATTATAT TACTTGGNAT GAATTATCCT ATGATCATCA 1160
     221 AGGTAAGGAA GTCTTGACTC CTAAGGCTTG GGACAGAAAT 1200
     223 GGGCAGGATT TNACGGCTCA CTTTACCACT AGTATTCCTT 1240
     225 TAAAAGGGAA TGTTCGTAAT CTCTCTGTCA AAATTAGAGA 1280
     227 GTGTACCGGG CTTGCNTGGG AATGGTGGCG TACGGTTTAT 1320
     229 GAAAAAACCG ATTTGCCACT AGTGCGTAAG CGGACGATTT 1360
     231 CTATTTGGGG AACAACTCTC TATCCNCAGG TAGANGATAA 1400
     233 GGTAGAAAAT GAC 1413
     237 (2) INFORMATION FOR SEQ ID NO: 3:
     239
              (i) SEQUENCE CHARACTERISTICS:
     240
                   (A) LENGTH: 471
     241
                   (B) TYPE: amino acid
     242
                   (C) STRANDEDNESS: unknown
     243
                   (D) TOPOLOGY: linear
W--> 245
             (ii) MOLECULE TYPE: amino acid
     247
             (vi) ORIGINAL SOURCE:
     248
                   (A) ORGANISM: S. pneumoniae
```

PATENT APPLICATION: US/10/785,673

DATE: 03/08/2004 TIME: 15:22:42

Input Set : A:\PTO.AMC.txt

25	Ω	(🗴	il s	EQUE:	NCE	DECC	DTDM	TON.	ano	T-0		_
		+ z.1	1, D	n I w		DESC	Y T L I	TON:	SEQ	TD.	NO:	3:
25	2 1	t AI	a AS	ш ту	S AL	a va	1 As	n As	p Ph			u Ala
			_	_		5				1	0	
25	4 Me1 -	t As:	n Ty	r Ası	o Ly	s Ly	s Ly	s Le	u Le	u Th	r Hi	s Gln
25.	5		1	5				2	0			
25	6 Gly	y Gl	u Se	r Ile	e Gl	u Ası	n Ar	a Ph	e T1	e Lv	s Gli	u Gly
25	7 25	5				3	0	J		Сцу	3!	
258	R Ast	n Gli	n T.e.	u Pro) 7 G			- 17-	7 77			-
25	3 1101	. 01.	1 116			יבט כ	ı Pin	e va		T TT	e GII	u Arg
			_	4(45			
260	У губ	з гъ	s Ar	g Sei	: Le	ı Sei	r Thi	r As	n Th:	r Se:	r Ası	o Ile
26:		50					5.					60
262	2 Ser	· Val	l Thi	r Ala	1 Thi	: Asr	ı Ası	o Se	r Ar	a Lei	ı Tvı	r Pro
263	3				65	5				7(
264	Glv	7 A1a	a T.e.	ı Lei			7) Gr	~ C1.	, Պե	, , , , , , , , , , , , , , , , , , ,	. .	ı Glu
265	<u>-</u>		75			- val	. Asi			тет	тес	ı Glu
		7			-	_		8)			
200	ASI	ASI	1 Pro	Thr	. Let	ı Let	ı Ala	a Vai	l Ası	o Arg	g Ala	a Pro
267						90					95	5
268	Met	Thr	туг :	Ser	· Ile	Asp	Lei	ı Pro	o Glv	z Lei	ı Ala	Ser
269)			100	ı	-			105			
270	Ser	Asn	Ser	Phe	T.e.11	Glr	1751	Cl,	100	, Dan e		Asn
271		110	1		LCU	. О11			ı ASL	PIC	ser	
272				70	<i>a</i> 1		115					120
	ser	ser	vai	. Arg	GTA	Ala	. Val	. Asr	ı Asp) Let	ı Leu	Ala
273					125					130)	
274	Lys	Trp	His	Gln	Asp	Tyr	Gly	Glr	ı Val	Asn	Asn	Val
275			135		_	_	_	140				
276	Pro	Ala	Ara	Met	Gln	Tur	G111	Tyro	т 7 о	mh.	. 77 -	77.7
277	145		9	1100	0111	150	GIU	г пуз	TIE	TUL		
278		Mot	C1	C11	Ŧ			_			155	
279	DEI	Met	Gru	Gln	ьeu	rÀs	val	Lys			Ser	Asp
				160					165			
281	Phe	Glu	Lys	Thr	Gly	Asn	Ser	Leu	Asp	Ile	Asp	Phe
282		170					175					180
283	Asn	Ser	Val	His	Ser	Ğlv	G] 11	Lvs	Gln	Tle	Gln	Ile
284					185	2		1370	0111	190	GLII	116
285	Val	Asn	Phe	Lys		Tlo	П	m	m1	190	_	
286		11011	195	цуз	GIII	тте	TÀT	lyr	Thr	val	Ser	Val
287	7	70.7		_	_			200				
	ASP	Α⊥а	vaı	Lys	Asn	Pro	Gly	Asp	Val	Phe	Gln	Asp
288	205					210					215	
289	Thr	Val	Thr	Val	Glu	Asp	Leu	Lys	Gln	Ara	Gly	Tle
290				220		•		4 -	225	9	<u>- 7</u>	110
291	Ser	Ala	Glu	Arg	Pro	T.011	Val	Фъг∽		0	Ser	** 1
292		230		9	110	пец		TÀT	тте	ser	ser	
	71-		C1	70	~ 3		235					240
200	Ата	1 AT.	сту	Arg	GIn	Val	Tyr	Leu	Lys	Leu	Glu	Thr
294					245					250		
295	Thr	Ser	Lys	Ser	Asp	Glu	Val	Glu	Ala	Ala	Phe	Glu
296			255					260				0.20
297	Ala	Leu	Tle	Lys	Glv	V=1	Lvc	1/2]	70.7 -	Dance	C1	mı
298	265	_			- L y	270	Ly S	val	TTG	LIO		rnr
299		Tra	T	C1-	T 1	2/0	70	_			275	
	GLU	ттБ	гуѕ	Gln	тте	ьeu	Asp	Asn		Glu	Val	Lys
300				280					285			

PATENT APPLICATION: US/10/785,673

DATE: 03/08/2004 TIME: 15:22:42

Input Set : A:\PTO.AMC.txt

```
301 Ala Val Ile Leu Gly Gly Asp Pro Ser Ser Gly Ala
              290
                                   295
       303 Arg Val Val Thr Gly Lys Val Asp Met Val Glu Asp
                          305
                                               310
      305 Leu Ile Gln Glu Gly Ser Arg Phe Thr Ala Asp His
               315
                                      320
      307 Pro Gly Leu Pro Ile Ser Tyr Thr Thr Ser Phe Leu
                              330
      309 Arg Asp Asn Val Val Ala Thr Phe Gln Asn Ser Thr
                      340
                                          345
      311 Asp Tyr Val Glu Thr Lys Val Thr Ala Tyr Arg Asn
      312 350
                                 355
      313 Gly Asp Leu Leu Leu Asp His Ser Gly Ala Tyr Val
                          365
                                              370
      315 Ala Gln Tyr Tyr Ile Thr Trp Asn Glu Leu Ser Tyr
      316 375
                                      380
      317 Asp His Gln Gly Lys Glu Val Leu Thr Pro Lys Ala
      318 385
                             390
      319 Trp Asp Arg Asn Gly Gln Asp Leu Thr Ala His Phe
                     400
                                          405
      321 Thr Thr Ser Ile Pro Leu Lys Gly Asn Val Arg Asn
      322
              410
                                 415
      323 Leu Ser Val Lys Ile Arg Glu Cys Thr Gly Leu Ala
      324
                        425
                                             430
     325 Trp Glu Trp Trp Arg Thr Val Tyr Glu Lys Thr Asp
      326
                 435
                                     440
     328 Leu Pro Leu Val Arg Lys Arg Thr Ile Ser Ile Trp
     329 445
                             450
     330 Gly Thr Thr Leu Tyr Pro Gln Val Glu Asp Lys Val
                     460
     332 Glu Asn Asp
             470
     336 (2) INFORMATION FOR SEQ ID NO: 4:
              (i) SEQUENCE CHARACTERISTICS:
     339
                   (A) LENGTH: 471
     340
                   (B) TYPE: amino acid
     341
                   (C) STRANDEDNESS: unknown
     342
                   (D) TOPOLOGY: linear
W--> 344
             (ii) MOLECULE TYPE: amino acid
     346
             (vi) ORIGINAL SOURCE:
     347
                   (A) ORGANISM: S. pneumoniae
     349
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
     351 Met Ala Asn Lys Ala Val Asn Asp Phe Ile Leu Ala
     352 1
                          5
W--> 353 Met Asn Tyr Asp Xaa Xaa Lys Leu Leu Thr His Gln
     354
                1.5
                                     20
    355 Gly Glu Ser Ile Glu Asn Arg Phe Xaa Lys Glu Gly
    356 25
                             30
    357 Asn Gln Leu Pro Xaa Glu Phe Val Xaa Xaa Glu Arg
```



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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/785,673

DATE: 03/05/2004 TIME: 12:01:54

Input Set : A:\17584036.txt

Output Set: N:\CRF4\03052004\J785673.raw

SEQUENCE LISTING

```
4 (1) GENERAL INFORMATION:
              (i) APPLICANT: MINETTI, CONCEICAO;
       7
                              MICHON, FRANCIS;
       8
                              PULLEN, JEFFREY K.;
       9
                              POLDVINO-BODNAR, MARYELLEN;
      10
                              LIANG, SHU-MEI;
      11
                              TAI, JOSEPH Y.
      13
             (ii) TITLE OF INVENTION: MODIFIED IMMUNOGENIC
      14
                                       PNEUMOLYSIN COMPOSITIONS AS VACCINES
     16
            (iii) NUMBER OF SEQUENCES: 18
             (iv) CORRESPONDENCE ADDRESS:
     18
     19
                   (A) ADDRESSEE: MORGAN & FINNEGAN, L.L.P.
     20
                   (B) STREET: 345 PARK AVENUE
     21
                   (C) CITY: NEW YORK
     22
                   (D) STATE: NEW YORK
     23
                   (E) COUNTRY: USA
     24
                   (F) ZIP: 10154
     26
             (v) COMPUTER READABLE FORM:
     27
                   (A) MEDIUM TYPE: FLOPPY DISK
     28
                   (B) COMPUTER: IBM PC COMPATIBLE
     29
                   (C) OPERATING SYSTEM: PC-DOS/MS-DOS
     30
                   (D) SOFTWARE: MICROSOFT WORD 97
     32
            (vi) CURRENT APPLICATION DATA:
C--> 33
                   (A) APPLICATION NUMBER: US/10/785,673
C--> 34
                   (B) FILING DATE: 23-Feb-2004
     35
                   (C) CLASSIFICATION:
     41
           (vii) PRIOR APPLICATION DATA:
     38
                   (A) APPLICATION NUMBER: 60/053,306
     39
                   (B) FILING DATE: 1997-07-21
     42
                  (A) APPLICATION NUMBER: 60/073,456
     43
                  (B) FILING DATE: 1998-02-02
     45
          (viii) ATTORNEY/AGENT INFORMATION:
     46
                  (A) NAME: DARRYL H. STEENSMA
     47
                  (B) REGISTRATION NUMBER: 43,155
     49
                  (C) REFERENCE/DOCKET NUMBER: 1758-4036US2
    51
            (ix) TELECOMMUNICATION INFORMATION:
    52
                  (A) TELEPHONE: (212) 758-4800
    53
                  (B) TELEFAX: (212) 751-6849
                  (C) TELEX: 421792
```

PATENT APPLICATION: US/10/785,673

DATE: 03/05/2004 TIME: 12:01:54

Input Set : A:\17584036.txt

Output Set: N:\CRF4\03052004\J785673.raw

ERRORED SEQUENCES

```
336 (2) INFORMATION FOR SEQ ID NO: 4:
               (i) SEQUENCE CHARACTERISTICS:
      339
                     (A) LENGTH: 471
      340
                     (B) TYPE: amino acid
      341
                     (C) STRANDEDNESS: unknown
      342
                     (D) TOPOLOGY: linear
   -> 344
              (ii) MOLECULE TYPE: amino acid
      346
              (vi) ORIGINAL SOURCE:
      347
                    (A) ORGANISM: S. pneumoniae
              (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
      349
      351 Met Ala Asn Lys Ala Val Asn Asp Phe Ile Leu Ala
                            5
                                               10
 W--> 353 Met Asn Tyr Asp Xaa Xaa Lys Leu Leu Thr His Gln
                   15
                                        20
      355 Gly Glu Ser Ile Glu Asn Arg Phe Xaa Lys Glu Gly
      356 25
                               30
      357 Asn Gln Leu Pro Xaa Glu Phe Val Xaa Xaa Glu Arg
      358
                       40
                                           45
      359 Lys Lys Arg Ser Leu Ser Thr Asn Thr Ser Asp Ile
      360 50
                                   55
     361 Xaa Val Xaa Ala Thr Xaa Asp Ser Arg Leu Tyr Pro
     362
                           65
                                               70
     363 Gly Ala Leu Leu Val Val Asp Glu Thr Xaa Leu Glu
               75
                                       80
     365 Asn Asn Pro Thr Leu Leu Ala Val Asp Arg Ala Pro
     366 85
                              90
     367 Met Thr Tyr Ser Xaa Xaa Leu Pro Gly Leu Ala Ser
     368
                     100
                                          105
     369 Ser Asp Ser Phe Leu Gln Val Glu Asp Pro Ser Asn
     370
           110
                                  115
     371 Ser Ser Val Arg Gly Ala Xaa Xaa Asp Leu Leu Ala
     372
                         125
                                              130
     373 Lys Trp His Gln Asp Tyr Gly Gln Val Asn Asn Val
                135
                                      140
     376 Pro Ala Arg Xaa Gln Tyr Glu Lys Xaa Thr Ala His
E--> 377 145
                             \$0 1<del>50</del>
                                                  ISS 155
     378 Ser Met Glu Gln Leu Lys Val Lys Phe Gly Ser Asp
E--> 379
                     160
                                          165
     380 Phe Glu Lys Xaa Gly Asn Ser Leu Asp Ile Asp Phe
E--> 381
             170
                                 175
     382 Asn Ser Val His Ser Gly Glu Lys Xaa Ile Gln Ile
E--> 383
                         185
                                              190
     384 Val Asn Xaa Lys Gln Ile Tyr Tyr Thr Val Ser Val
E--> 385
                 195
                                     200
     386 Asp Ala Val Lys Asn Pro Gly Asp Val Phe Gln Asp
E--> 387 205
                             210
     388 Thr Val Thr Val Glu Asp Leu Lys Gln Arg Gly Ile
E--> 389
                     220
```

PATENT APPLICATION: US/10/785,673

DATE: 03/05/2004 TIME: 12:01:54

Input Set : A:\17584036.txt

E>	390 39 :	0 Se: 1	r Ala 23 0	a Glu	ı Arç	y Pro	Le	u Val	1 Туз	r Ile	e Sei	: Xaa	a Val
					. 7\ ~~ a			23!	5	_			240
E>	> 39.	•				245	5				250)	ı Thr
ъ.	394	Thi	c Sei	Xaa	Ser	Xaa	ı Glı	ı Val	l Glı	ı Ala	Ala	Phe	e Glu
E>	, 39:)		255)				260	1			
F>	305	7 265	. тег	1 TTE	Lys	GLY	7 Val	L Lys	8 Val	. Ala	Pro	Glr	Thr
E>		200	,				2/()				275	:
E>	300) GIC	1 114	ь гуз	GIN	тте	Let	ı Asp) Asn			Val	Lys
<u> </u>			. 17-1	T1 -	280		~ 1	_	_	285			
E>	400	, ATO	290	тте	Leu	GIY	, GTZ	Asp	Pro	Ser	Ser	Gly	Ala
					Прх	C1	7	295	, ,				300
E>	403		, vai	Val	TIII	305	тЛг	val	. Asp	Met			Asp
			Tle	Gln	Glu	203 C1v	Sar	. 7	. Dh.	m1	310	_	His
E>	405	,		315	Oru	Gry	ser	Arg	320	Inr	Ата	Asp	His
	406	Pro	Glv			Tle	Ser	• Ттг	720 Thr	Th∽	Co.~	Dl	Leu
E>	407	325	-				330	- <u>y</u> -	1111	1111	ser	335	
	408	Arg	Asp	Asn	Val	Val	Ala	Thr	Phe	Gln	Asn	Ser	Thr
E>	409				340					345			
	410	Asp	Tyr	Val	Glu	Thr	Lys	Val	Thr	Ala	Tvr	Ara	Agn
E>	411		350					355					360
	412	Gly	Asp	Leu	Leu	Leu	Asp	His	Ser	Gly	Ala	Tvr	Val
E>	413					365					370		
_	414	Ala	Gln	Tyr	Tyr	Ile	Thr	Trp	Xaa	Glu	Leu	Ser	Tyr
E>	415			375					380				
	416	Asp	His	Gln	Gly	Lys	Glu	Val	Leu	Thr	Pro	Lys	Ala
E>	41 /	303					390					305	
E>	410	Trp	Asp	Arg	Asn	Gly	Gln	Asp	Leu	Thr	Ala	His	Phe
E>		Thr	mb ×	Com	400	Б	_	_		405			
E>	421	* 111	410	ser	тте	Pro	Leu	Lys	GTY	Asn	Val	Arg	Asn
		T.e.11		17 a 1	Two	Tlo	7\	415	•	-		_	420
E>	424	11Cu	DCI	vai	пуз	425	Arg	GIU	Cys	Thr		Leu	Ala
		Trp	G] 11	Trn			ጥኮሎ	T/~ 1	П	Glu	430	m.i	_
E>	426			435	ı⊥p	111 g	1111		440	GIU	ьуs	Thr	Asp
	427	Leu			Val	Ara	T.378	Δκα	Thr	Ile	C	- 1 -	m
E>	428	445				9	450	11 1 9	1111	116		455	Trp
	429	Gly	Thr	Thr	Leu	Tvr	Pro	Gln	Val	Glu .	Δen	435 Tuc	V-1
E>	430	_			460	<i></i>				465	. Tab	пÌЭ	val
	431	Glu	Asn .	Asp									
E>	432		470	=									

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/785,673

DATE: 03/05/2004 TIME: 12:01:55

Input Set : A:\17584036.txt

Output Set: N:\CRF4\03052004\J785673.raw

L:6 M:220 C: Keyword misspelled or invalid format, [(i) APPLICANT:] L:33 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:] L:34 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:] L:72 M:111 C: (47) String data converted to upper case, M:111 Repeated in SeqNo=1 L:162 M:111 C: (47) String data converted to upper case, M:111 Repeated in SeqNo=2 L:245 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=3 L:344 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=4 L:353 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:12 M:341 Repeated in SeqNo=4 L:377 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:4 M:332 Repeated in SeqNo=4 L:443 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=5 L:450 M:111 C: (47) String data converted to upper case, L:461 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=6 L:468 M:111 C: (47) String data converted to upper case, L:480 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=7 L:487 M:111 C: (47) String data converted to upper case, L:498 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=8 L:505 M:111 C: (47) String data converted to upper case, L:517 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=9 L:524 M:111 C: (47) String data converted to upper case, L:535 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=10 L:542 M:111 C: (47) String data converted to upper case, L:553 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=11 L:560 M:111 C: (47) String data converted to upper case, L:572 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=12 L:579 M:111 C: (47) String data converted to upper case, L:590 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=13 L:597 M:111 C: (47) String data converted to upper case, L:609 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=14 L:616 M:111 C: (47) String data converted to upper case, L:626 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=15 L:633 M:111 C: (47) String data converted to upper case, L:644 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=16 L:651 M:111 C: (47) String data converted to upper case, L:663 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=17 L:670 M:111 C: (47) String data converted to upper case, L:681 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=18 L:688 M:111 C: (47) String data converted to upper case,